11333US.ST25.txt
SEQUENCE LISTING

<110>   Siegfried Burggraf

<120>   Method for Detecting Nucleic Acids with internal Control of the
        Amplification

<130>   11333US

<150>   EP 03015027.0
<151>   2003-07-02

<150>   PCT/EP04/02680
<151>   2004-03-15

<160>   2

<170>   PatentIn version 3.1

<210>   1
<211>   188
<212>   DNA
<213>   Hepatitis B virus

<220>
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<222>   (1)..(188)
<223>   GenX

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cttggggctc tacgccctct ttgccgtctg ccgttccagc caaccacggg gcgcacctct   120

ctttacgcgg tctccccgtc tgtgccttct catctgccgg tccgtgtgca cttcgcttca   180

cctctgca   188

<210>   2
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<212>   DNA
<213>   Artificial sequence

<220>
<223>   This sequence is an artificial one which only partially coincides
        with GenX of Hepatitis B (SEQ ID NO:1)

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<223>   This sequence is an artificial one which only partially coincides
        with GenX of Hepatitis B (SEQ ID NO:1)

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ttacgcggtc taccagtcta tgccttatca tctgcctgtg cacttcgctt cacctctgca   120

Applicant

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<110> ApplicantName : Siegfried Burggraf

Application Project

<120> Title : Method for Detecting Nucleic Acids with Internal Control of the
        Amplification
<130> AppFileReference : 11333US
<140> CurrentAppNumber :
<141> CurrentFilingDate : ____-__-__

Earlier Applications

<150> PriorAppNumber : EP 03015027.0
<151> PriorFilingDate : 2003-07-03

<150> PriorAppNumber : PCT/EP04/02680
<151> PriorFilingDate : 2004-03-15

Sequence

<213> OrganismName : Hepatitis B virus
<400> PreSequenceString :
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cttggggctc tacgccctct ttgccgtctg ccgttccagc caaccacggg gcgcacctct   120
ctttacgcgg tctccccgtc tgtgccttct catctgccgg tccgtgtgca cttcgcttca   180
cctctgca   188
<212> Type : DNA
<211> Length : 188
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        SequenceDescription :

Feature

Sequence: SEQ ID NO:1:
<221> FeatureKey : source
<222> LocationFrom : 1
<222> LocationTo : 188
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        CDSJoin : No

Sequence

<213> OrganismName : Artificial sequence
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<212> Type : DNA
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Feature

Sequence: SEQ ID NO:2:
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Other Information : This sequence is an artificial one which only partially coincides with GenX of Hepatitis B (SEQ ID NO:1)
CDSJoin : No